

**Amendments to the Claims**

Please cancel claim 5 and amend claims 1, 6, 7, 16, 25 and 27 under the provisions of 37 C.F.R. §1.121, as set forth in the Federal Register on June 30, 2003, as follows:

1. (Currently Amended) A pharmaceutical composition comprising:  
an aqueous carrier;  
from 0.1 mg/ml to 20 mg/ml of the composition of a pharmaceutically acceptable salt of
  - a) a peptide comprising at least 12 and at most 30 consecutive amino acids having a sequence corresponding to
    - (i) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a human monoclonal anti-DNA 16/6 Id antibody, or
    - (ii) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a pathogenic anti-DNA monoclonal antibody that induces a systemic lupus erythematosus (SLE)-like disease response in mice, or
  - b) a peptide comprising consecutive amino acids having the sequence
    - (i) TGYX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>X<sub>4</sub>X<sub>5</sub>QSPEKSLEWIG (SEQ ID NO:11)  
wherein X<sub>1</sub> is Met, Ala or Val; X<sub>2</sub> is Gln, Asp, Glu or Arg; X<sub>3</sub> is Trp or Ala; X<sub>4</sub> is Val or Ser; and X<sub>5</sub> is Lys, Glu or Ala;
    - (ii) EINPSTGGX<sub>6</sub>X<sub>7</sub>X<sub>8</sub>X<sub>9</sub>X<sub>10</sub>X<sub>11</sub>X<sub>12</sub>KAKAT (SEQ ID NO:12)  
wherein X<sub>6</sub> and X<sub>7</sub> are each Thr, Val or Ala; X<sub>8</sub>

is Tyr or Phe; X<sub>9</sub> is Asn or Asp; X<sub>10</sub> is Gln or Glu; X<sub>11</sub> is Lys or Glu, and X<sub>12</sub> is Phe or Tyr;

(iii) YYCARX<sub>13</sub>X<sub>14</sub>X<sub>15</sub>X<sub>16</sub>PYAX<sub>17</sub>X<sub>18</sub>YWGQGS (SEQ ID NO:13)  
wherein X<sub>13</sub> is Phe, Thr or Gly; X<sub>14</sub> is Leu, Ala or Ser; X<sub>15</sub> is Trp or Ala; X<sub>16</sub> is Glu or Lys; X<sub>17</sub> is Met or Ala, and X<sub>18</sub> is Asp, Lys or Ser;

(iv) GYNX<sub>19</sub>X<sub>20</sub>X<sub>21</sub>X<sub>22</sub>X<sub>23</sub>X<sub>24</sub>SHGX<sub>25</sub>X<sub>26</sub>LEWIG (SEQ ID NO:14)  
wherein X<sub>19</sub> is Met or Ala; X<sub>20</sub> is Asn, Asp or Arg; X<sub>21</sub> is Trp or Ala; X<sub>22</sub> is Val or Ser; X<sub>23</sub> is Lys or Glu; X<sub>24</sub> is Gln or Ala; X<sub>25</sub> is Lys or Glu, and X<sub>26</sub> is Ser or Ala;

(v) YYCARX<sub>27</sub>X<sub>28</sub>X<sub>29</sub>YGX<sub>30</sub>X<sub>31</sub>X<sub>32</sub>GQTL (SEQ ID NO:15)  
wherein X<sub>27</sub> is Ser or Phe; X<sub>28</sub> is Gly or Ala; X<sub>29</sub> is Arg, Ala or Glu; X<sub>30</sub> is Asn or Asp; X<sub>31</sub> is Tyr or Phe, and X<sub>32</sub> is Trp, His or Ala;

(vi) X<sub>33</sub>YYWSWIX<sub>34</sub>QX<sub>35</sub>PX<sub>36</sub>X<sub>37</sub>GX<sub>38</sub>EWIG (SEQ ID NO:16)  
wherein X<sub>33</sub> is Gly or Thr Gly; X<sub>34</sub> is Arg or Lys; X<sub>35</sub> is Pro or Ser; X<sub>36</sub> is Gly or Glu; X<sub>37</sub> is Lys or Asp; and X<sub>38</sub> is Glu, Leu or Ser;

(vii) YYCARX<sub>39</sub>LLX<sub>40</sub>X<sub>41</sub>X<sub>42</sub>X<sub>43</sub>X<sub>44</sub>DVDYX<sub>45</sub>GX<sub>46</sub>DV (SEQ ID NO:17)

wherein X<sub>39</sub> is Gly or Phe; X<sub>40</sub> is Arg or Ala; X<sub>41</sub> is Gly or Ala; X<sub>42</sub> is Gly or Ala; X<sub>43</sub> is Trp or Ala; X<sub>44</sub> is Asn or Ala; X<sub>45</sub> is Tyr or Trp; X<sub>46</sub> is Met or Gln;

(viii) FSGYYWS (SEQ ID NO:8);

(ix) EINHSGSTNYKTSLS (SEQ ID NO:9); or

(x) GLLRGGWNDVDYYYGMDV (SEQ ID NO:10), or

c) a peptide comprising consecutive amino acids having a sequence of any of a) and b), or at least two of the sequences in (a)(i), (a)(ii) and (b)(i) through

(b) (x), or

- d) a peptide comprising consecutive amino acids having a sequence comprising at least two identical sequences included in (a)(i), (a)(ii) and (b)(i) through (b)(x); and

~~a solubility enhancer selected from the group consisting of dimethyl acetamide, polyethylene glycol, polyoxylated easter oil, N-methyl-2-pyrrolidinone, 1-ethenyl-2-pyrrolidinone, polyoxyethylene sorbitan esters, and a substituted  $\beta$ -cyclodextrin,~~

wherein both the peptide and ~~the solubility enhancer~~  
the substituted  $\beta$ -cyclodextrin are dissolved in the aqueous carrier; and

wherein the pharmaceutical composition has a pH between 4 and 9.

2. (Original) The pharmaceutical composition of claim 1, wherein at least 0.5 mg/ml of the composition is the pharmaceutically acceptable salt of the peptide.
3. (Previously Presented) The pharmaceutical composition of claim 1, wherein the peptide has a sequence selected from the group consisting of:

NH<sub>2</sub>- Thr Gly Tyr Tyr Met Gln Trp Val Lys Gln Ser Pro Glu Lys  
Ser Leu Glu-Trp Ile Gly-COOH (SEQ ID NO:1);

NH<sub>2</sub>- Glu Ile Asn Pro Ser Thr Gly Gly Thr Thr Tyr Asn Gln Lys  
Phe Lys Ala Lys Ala Thr-COOH (SEQ ID NO:2);

NH<sub>2</sub>- Tyr Tyr Cys Ala Arg Phe Leu Trp Glu Pro Tyr Ala Met Asp  
Tyr Trp Gly Gln Gly Ser-COOH (SEQ ID NO:3);

NH<sub>2</sub>- Gly Tyr Asn Met Asn Trp Val Lys Gln Ser His Gly Lys Ser  
Leu Glu Trp Ile Gly-COOH (SEQ ID NO:4);

NH<sub>2</sub>- Tyr Tyr Cys Ala Arg Ser Gly Arg Tyr Gly Asn Tyr Trp Gly  
Gln Thr Leu -COOH (SEQ ID NO:5);

NH<sub>2</sub>-Gly Tyr Tyr Trp Ser Trp Ile Arg Gln Pro Pro Gly Lys Gly  
Glu Glu Trp Ile Gly-COOH (SEQ ID NO:6);

NH<sub>2</sub>-Tyr Tyr Cys Ala Arg Gly Leu Leu Arg Gly Gly Trp Asn Asp  
Val Asp Tyr Tyr Gly Met Asp Val-COOH (SEQ ID NO:7);

NH<sub>2</sub>- Phe Ser Gly Tyr Tyr Trp Ser-COOH (SEQ ID NO:8);

NH<sub>2</sub>- Glu Ile Asn His Ser Gly Ser Thr Asn Tyr Lys Thr Ser Leu  
Lys Ser-COOH (SEQ ID NO:9); and

NH<sub>2</sub>- Gly Leu Leu Arg Gly Gly Trp Asn Asp Val Asp Tyr Tyr  
Tyr Gly Met Asp Val-COOH (SEQ ID NO:10).

4. (Original) The pharmaceutical composition of claim 1, wherein the peptide comprises consecutive amino acids having the sequence

X<sub>33</sub>YYWSWIX<sub>34</sub>QX<sub>35</sub>PX<sub>36</sub>X<sub>37</sub>GX<sub>38</sub>EWIG (SEQ ID NO:16)

wherein X<sub>33</sub> is Gly or Thr Gly; X<sub>34</sub> is Arg or Lys; X<sub>35</sub> is Pro or Ser; X<sub>36</sub> is Gly or Glu; X<sub>37</sub> is Lys or Asp; and X<sub>38</sub> is Glu, Leu or Ser.

5. (Canceled).

6. (Currently Amended) The pharmaceutical composition of claim—51, wherein the substituted  $\beta$ -cyclodextrin is a hydroxypropyl, a sulfobutyl ether, or asulfopropyl ether substituted  $\beta$ -cyclodextrin.

7. (Currently Amended) The pharmaceutical composition of claim 6, wherein the substituted  $\beta$ -cyclodextrin is a

~~substituted~~-sulfobutyl ether substituted  $\beta$ -cyclodextrin.

8. (Previously Presented) The pharmaceutical composition of claim 1, wherein the concentration of peptide in solution is at least 1 mg/ml.
9. (Previously Presented) The pharmaceutical composition of claim 1, wherein the concentration of peptide in solution is at least 2.5 mg/ml.
10. (Previously Presented) The pharmaceutical composition of claim 1, wherein the composition has a pH between 6.5 and 8.5.
11. (Previously Presented) The pharmaceutical composition of claim 10, wherein the pharmaceutical composition has a pH between 7.5 and 8.5.
12. (Previously Presented) The pharmaceutical composition of claim 1, wherein the pharmaceutically acceptable salt is an acetate salt.
13. (Original) The pharmaceutical composition of claim 5, wherein the pharmaceutically acceptable salt is an acetate salt, and the substituted  $\beta$ -cyclodextrin is hepta-(sulfobutyl ether)- $\beta$ -cyclodextrin.
14. (Withdrawn) A method of alleviating symptoms of systemic lupus erythematosus (SLE) in a human subject comprising administering to the human subject the pharmaceutical composition of claim 1 in an amount effective to

alleviate the symptoms of the SLE in the human subject.

15. (Canceled)

16. (Currently Amended) A process for manufacturing the pharmaceutical composition of claim 1, comprising the steps of:

a) preparing a solution of ~~dimethyl acetamide, polyethylene glycol, polyoxylated castor oil, N methyl 2-pyrrolidinone, 1 ethenyl 2-pyrrolidinone, polyoxyethylene sorbitan esters, or a~~ substituted  $\beta$ -cyclodextrin in an aqueous carrier;

b) adding a pharmaceutically acceptable salt of

1) a peptide comprising at least 12 and at most 30 consecutive amino acids having a sequence corresponding to

(i) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a human monoclonal anti-DNA 16/6 Id antibody, or

(ii) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a pathogenic anti-DNA monoclonal antibody that induces a systemic lupus erythematosus (SLE)-like disease response in mice,

2) a peptide comprising amino acids having the sequence

(i) TGYX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>X<sub>4</sub>X<sub>5</sub>QSPEKSLEWIG (SEQ ID NO:11)

wherein X<sub>1</sub> is Met, Ala or Val; X<sub>2</sub> is Gln, Asp,

Glu or Arg; X<sub>3</sub> is Trp or Ala; X<sub>4</sub> is Val or Ser;  
and X<sub>5</sub> is Lys, Glu or Ala;

- (ii) EINPSTGGX<sub>6</sub>X<sub>7</sub>X<sub>8</sub>X<sub>9</sub>X<sub>10</sub>X<sub>11</sub>X<sub>12</sub>KAKAT (SEQ ID NO:12)  
wherein X<sub>6</sub> and X<sub>7</sub> are each Thr, Val or Ala; X<sub>8</sub>  
is Tyr or Phe; X<sub>9</sub> is Asn or Asp; X<sub>10</sub> is Gln or  
Glu; X<sub>11</sub> is Lys or Glu, and X<sub>12</sub> is Phe or Tyr;
- (iii) YYCARX<sub>13</sub>X<sub>14</sub>X<sub>15</sub>X<sub>16</sub>PYAX<sub>17</sub>X<sub>18</sub>YWGQGS (SEQ ID NO:13)  
wherein X<sub>13</sub> is Phe, Thr or Gly; X<sub>14</sub> is Leu, Ala  
or Ser; X<sub>15</sub> is Trp or Ala; X<sub>16</sub> is Glu or Lys; X<sub>17</sub>  
is Met or Ala, and X<sub>18</sub> is Asp, Lys or Ser;
- (iv) GYNX<sub>19</sub>X<sub>20</sub>X<sub>21</sub>X<sub>22</sub>X<sub>23</sub>X<sub>24</sub>SHGX<sub>25</sub>X<sub>26</sub>LEWIG (SEQ ID NO:14)  
wherein X<sub>19</sub> is Met or Ala; X<sub>20</sub> is Asn, Asp or  
Arg; X<sub>21</sub> is Trp or Ala; X<sub>22</sub> is Val or Ser; X<sub>23</sub> is  
Lys or Glu; X<sub>24</sub> is Gln or Ala; X<sub>25</sub> is Lys or Glu,  
and X<sub>26</sub> is Ser or Ala;
- (v) YYCARX<sub>27</sub>X<sub>28</sub>X<sub>29</sub>YGX<sub>30</sub>X<sub>31</sub>X<sub>32</sub>GQTL (SEQ ID NO:15)  
wherein X<sub>27</sub> is Ser or Phe; X<sub>28</sub> is Gly or Ala; X<sub>29</sub>  
is Arg, Ala or Glu; X<sub>30</sub> is Asn or Asp; X<sub>31</sub> is Tyr  
or Phe, and X<sub>32</sub> is Trp, His or Ala;
- (vi) X<sub>33</sub>YYWSWIX<sub>34</sub>QX<sub>35</sub>PX<sub>36</sub>X<sub>37</sub>GX<sub>38</sub>EWIG (SEQ ID NO:16)  
wherein X<sub>33</sub> is Gly or Thr Gly; X<sub>34</sub> is Arg or Lys;  
X<sub>35</sub> is Pro or Ser; X<sub>36</sub> is Gly or Glu; X<sub>37</sub> is Lys  
or Asp; and X<sub>38</sub> is Glu, Leu or Ser;
- (vii) YYCARX<sub>39</sub>LLX<sub>40</sub>X<sub>41</sub>X<sub>42</sub>X<sub>43</sub>X<sub>44</sub>DVDYX<sub>45</sub>GX<sub>46</sub>DV (SEQ ID NO:17)  
wherein X<sub>39</sub> is Gly or Phe; X<sub>40</sub> is Arg or Ala; X<sub>41</sub>  
is Gly or Ala; X<sub>42</sub> is Gly or Ala; X<sub>43</sub> is Trp or  
Ala; X<sub>44</sub> is Asn or Ala; X<sub>45</sub> is Tyr or Trp; X<sub>46</sub> is  
Met or Gln;
- (viii) FSGYYWS (SEQ ID NO:8);
- (ix) EINHSGSTNYKTSLS (SEQ ID NO:9); or
- (x) GLLRGGWNDVDYYYGMDV (SEQ ID NO:10), or

- 3) a peptide comprising consecutive amino acids having a sequence of any of a) and b), or at least two of the sequences in (a)(i), (a)(ii) and (b)(i) through (b)(x), or
- 4) a peptide comprising consecutive amino acids having a sequence comprising at least two identical sequences included in (a)(i), (a)(ii) and (b)(i) through (b)(x);
- c) adjusting the pH of the solution of step b) until the peptide dissolves in the solution; and
- d) if necessary, adjusting the pH of the solution of step c) to a pH of 4-9, thereby manufacturing the pharmaceutical composition.

Claims 17-23. (Canceled)

- 24. (Previously Presented) A composition prepared by the process of claim 16.
- 25. (Currently Amended) A lyophilized pharmaceutical composition comprising from 0.1 mg/ml to 20 mg/ml of the composition of a pharmaceutically acceptable salt of
  - a) a peptide comprising at least 12 and at most 30 consecutive amino acids having a sequence corresponding to
    - (i) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a human monoclonal anti-DNA 16/6 Id antibody, or
    - (ii) a sequence of amino acids found within a complementarity-determining region (CDR) of a



heavy or a light chain of a pathogenic anti-DNA monoclonal antibody that induces a systemic lupus erythematosus (SLE)-like disease response in mice, or

b) a peptide comprising consecutive amino acids having the sequence

(i) TGYX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>X<sub>4</sub>X<sub>5</sub>QSPEKSLEWIG (SEQ ID NO:11)

wherein X<sub>1</sub> is Met, Ala or Val; X<sub>2</sub> is Gln, Asp, Glu or Arg; X<sub>3</sub> is Trp or Ala; X<sub>4</sub> is Val or Ser; and X<sub>5</sub> is Lys, Glu or Ala;

(ii) EINPSTGGX<sub>6</sub>X<sub>7</sub>X<sub>8</sub>X<sub>9</sub>X<sub>10</sub>X<sub>11</sub>X<sub>12</sub>KAKAT (SEQ ID NO:12)

wherein X<sub>6</sub> and X<sub>7</sub> are each Thr, Val or Ala; X<sub>8</sub> is Tyr or Phe; X<sub>9</sub> is Asn or Asp; X<sub>10</sub> is Gln or Glu; X<sub>11</sub> is Lys or Glu, and X<sub>12</sub> is Phe or Tyr;

(iii) YYCARX<sub>13</sub>X<sub>14</sub>X<sub>15</sub>X<sub>16</sub>PYAX<sub>17</sub>X<sub>18</sub>YWGQGS (SEQ ID NO:13)

wherein X<sub>13</sub> is Phe, Thr or Gly; X<sub>14</sub> is Leu, Ala or Ser; X<sub>15</sub> is Trp or Ala; X<sub>16</sub> is Glu or Lys; X<sub>17</sub> is Met or Ala, and X<sub>18</sub> is Asp, Lys or Ser;

(iv) GYNX<sub>19</sub>X<sub>20</sub>X<sub>21</sub>X<sub>22</sub>X<sub>23</sub>X<sub>24</sub>SHGX<sub>25</sub>X<sub>26</sub>LEWIG (SEQ ID NO:14)

wherein X<sub>19</sub> is Met or Ala; X<sub>20</sub> is Asn, Asp or Arg; X<sub>21</sub> is Trp or Ala; X<sub>22</sub> is Val or Ser; X<sub>23</sub> is Lys or Glu; X<sub>24</sub> is Gln or Ala; X<sub>25</sub> is Lys or Glu, and X<sub>26</sub> is Ser or Ala;

(v) YYCARX<sub>27</sub>X<sub>28</sub>X<sub>29</sub>YGX<sub>30</sub>X<sub>31</sub>X<sub>32</sub>GQTL (SEQ ID NO:15)

wherein X<sub>27</sub> is Ser or Phe; X<sub>28</sub> is Gly or Ala; X<sub>29</sub> is Arg, Ala or Glu; X<sub>30</sub> is Asn or Asp; X<sub>31</sub> is Tyr or Phe, and X<sub>32</sub> is Trp, His or Ala;

(vi) X<sub>33</sub>YYWSWIX<sub>34</sub>QX<sub>35</sub>PX<sub>36</sub>X<sub>37</sub>GX<sub>38</sub>EWIG (SEQ ID NO:16)

wherein X<sub>33</sub> is Gly or Thr Gly; X<sub>34</sub> is Arg or Lys; X<sub>35</sub> is Pro or Ser; X<sub>36</sub> is Gly or Glu; X<sub>37</sub> is Lys or Asp; and X<sub>38</sub> is Glu, Leu or Ser;

(vii) YYCARX<sub>39</sub>LLX<sub>40</sub>X<sub>41</sub>X<sub>42</sub>X<sub>43</sub>X<sub>44</sub>DVDYX<sub>45</sub>GX<sub>46</sub>DV (SEQ ID NO:17)

wherein X<sub>39</sub> is Gly or Phe; X<sub>40</sub> is Arg or Ala; X<sub>41</sub> is Gly or Ala; X<sub>42</sub> is Gly or Ala; X<sub>43</sub> is Trp or Ala; X<sub>44</sub> is Asn or Ala; X<sub>45</sub> is Tyr or Trp; X<sub>46</sub> is Met or Gln;

(viii) FSGYYWS (SEQ ID NO:8);

(ix) EINHSGSTNYKTSLS (SEQ ID NO:9); or

(x) GLLRGGWNDVDYYYGMDV (SEQ ID NO:10), or

c) a peptide comprising consecutive amino acids having a sequence of any of a) and b), or at least two of the sequences in (a)(i), (a)(ii) and (b)(i) through (b)(x), or

d) a peptide comprising consecutive amino acids having a sequence comprising at least two identical sequences included in (a)(i), (a)(ii) and (b)(i) through (b)(x); and

~~a solubility enhancer selected from the group consisting of dimethyl acetamide, polyethylene glycol, polyoxylated castor oil, N-methyl-2-pyrrolidinone, 1-ethenyl-2-pyrrolidinone, polyoxyethylene sorbitan esters, and a substituted  $\beta$ -cyclodextrin.~~

26. (Canceled)

27. (Currently Amended) A process for manufacturing the lyophilized pharmaceutical composition of claim 25, comprising the steps of:

a) preparing a solution of ~~dimethyl acetamide, polyethylene glycol, polyoxylated castor oil, N-methyl-2-pyrrolidinone, 1-ethenyl-2-~~

~~pyrrolidinone, polyoxyethylene sorbitan esters, or a~~  
substituted  $\beta$ -cyclodextrin in an aqueous carrier;

b) adding a pharmaceutically acceptable salt of

1) a peptide comprising at least 12 and at most 30 consecutive amino acids having a sequence corresponding to

(i) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a human monoclonal anti-DNA 16/6 Id antibody, or

(ii) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a pathogenic anti-DNA monoclonal antibody that induces a systemic lupus erythematosus (SLE)-like disease response in mice,

2) a peptide comprising amino acids having the sequence

(i) TGYX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>X<sub>4</sub>X<sub>5</sub>QSPEKSLEWIG (SEQ ID NO:11)

wherein X<sub>1</sub> is Met, Ala or Val; X<sub>2</sub> is Gln, Asp, Glu or Arg; X<sub>3</sub> is Trp or Ala; X<sub>4</sub> is Val or Ser; and X<sub>5</sub> is Lys, Glu or Ala;

(ii) EINPSTGGX<sub>6</sub>X<sub>7</sub>X<sub>8</sub>X<sub>9</sub>X<sub>10</sub>X<sub>11</sub>X<sub>12</sub>KAKAT (SEQ ID NO:12)

wherein X<sub>6</sub> and X<sub>7</sub> are each Thr, Val or Ala; X<sub>8</sub> is Tyr or Phe; X<sub>9</sub> is Asn or Asp; X<sub>10</sub> is Gln or Glu; X<sub>11</sub> is Lys or Glu, and X<sub>12</sub> is Phe or Tyr;

(iii) YYCARX<sub>13</sub>X<sub>14</sub>X<sub>15</sub>X<sub>16</sub>PYAX<sub>17</sub>X<sub>18</sub>YWGQGS (SEQ ID NO:13)

wherein X<sub>13</sub> is Phe, Thr or Gly; X<sub>14</sub> is Leu, Ala or Ser; X<sub>15</sub> is Trp or Ala; X<sub>16</sub> is Glu or Lys; X<sub>17</sub> is Met or Ala, and X<sub>18</sub> is Asp, Lys or Ser;

(iv) GYNX<sub>19</sub>X<sub>20</sub>X<sub>21</sub>X<sub>22</sub>X<sub>23</sub>X<sub>24</sub>SHGX<sub>25</sub>X<sub>26</sub>LEWIG (SEQ ID NO:14)

wherein  $X_{19}$  is Met or Ala;  $X_{20}$  is Asn, Asp or Arg;  $X_{21}$  is Trp or Ala;  $X_{22}$  is Val or Ser;  $X_{23}$  is Lys or Glu;  $X_{24}$  is Gln or Ala;  $X_{25}$  is Lys or Glu, and  $X_{26}$  is Ser or Ala;

(v) YYCARX<sub>27</sub>X<sub>28</sub>X<sub>29</sub>YGX<sub>30</sub>X<sub>31</sub>X<sub>32</sub>GQTL (SEQ ID NO:15)

wherein  $X_{27}$  is Ser or Phe;  $X_{28}$  is Gly or Ala;  $X_{29}$  is Arg, Ala or Glu;  $X_{30}$  is Asn or Asp;  $X_{31}$  is Tyr or Phe, and  $X_{32}$  is Trp, His or Ala;

(vi) X<sub>33</sub>YYWSWIX<sub>34</sub>QX<sub>35</sub>PX<sub>36</sub>X<sub>37</sub>GX<sub>38</sub>EWIG (SEQ ID NO:16)

wherein  $X_{33}$  is Gly or Thr Gly;  $X_{34}$  is Arg or Lys;  $X_{35}$  is Pro or Ser;  $X_{36}$  is Gly or Glu;  $X_{37}$  is Lys or Asp; and  $X_{38}$  is Glu, Leu or Ser;

(vii) YYCARX<sub>39</sub>LLX<sub>40</sub>X<sub>41</sub>X<sub>42</sub>X<sub>43</sub>X<sub>44</sub>DVDYX<sub>45</sub>GX<sub>46</sub>DV (SEQ ID NO:17)

wherein  $X_{39}$  is Gly or Phe;  $X_{40}$  is Arg or Ala;  $X_{41}$  is Gly or Ala;  $X_{42}$  is Gly or Ala;  $X_{43}$  is Trp or Ala;  $X_{44}$  is Asn or Ala;  $X_{45}$  is Tyr or Trp;  $X_{46}$  is Met or Gln;

(viii) FSGYYWS (SEQ ID NO:8);

(ix) EINHSGSTNYKTSLS (SEQ ID NO:9); or

(x) GLLRGGWNDVDYYYGMDV (SEQ ID NO:10), or

3) a peptide comprising consecutive amino acids having a sequence of any of a) and b), or at least two of the sequences in (a)(i), (a)(ii) and (b)(i) through (b)(x), or

4) a peptide comprising consecutive amino acids having a sequence comprising at least two identical sequences included in (a)(i), (a)(ii) and (b)(i) through (b)(x);

c) adjusting the pH of the solution of step b) until the peptide dissolves in the solution;

- d) if necessary, adjusting the pH of the solution of step c) to a pH of 4-9, thereby manufacturing the pharmaceutical composition; and
- e) lyophilizing the pharmaceutical composition of step d) by:
  - a-i) lowering the temperature of the pharmaceutical composition to  $-40^{\circ}\text{C}$ ;
  - a-ii) holding the temperature at  $-40^{\circ}\text{C}$  for a predetermined time;
  - a-iii) raising the temperature of the solution to  $20^{\circ}\text{C}$ ;
  - a-iv) holding the temperature at  $20^{\circ}\text{C}$  for a predetermined time; and
  - a-v) reducing the pressure and holding the temperature at  $20^{\circ}\text{C}$  for a predetermined time, thereby lyophilizing the pharmaceutical composition;or
  - b-i) lowering the temperature of the pharmaceutical composition to  $-45^{\circ}\text{C}$ ;
  - b-ii) holding the temperature at  $-45^{\circ}\text{C}$  for a predetermined time;
  - b-iii) raising the temperature of the solution to  $-20^{\circ}\text{C}$ ;
  - b-iv) raising the temperature of the solution to  $25^{\circ}\text{C}$ ;and
  - b-v) holding the temperature at  $25^{\circ}\text{C}$  for a predetermined time, thereby lyophilizing the pharmaceutical composition.

Claims 28-35. (Canceled)

36. (Previously Presented) The process of claim 27, wherein

step a-i) is performed within 2 hours;  
step a-ii) is performed within 3 hours;  
step a-iii) is performed over 13 hours and at a  
pressure of 110 $\mu$ bar;  
step a-iv) is performed over 13 hours and at a  
pressure of 110 $\mu$ bar; and  
step a-v) is performed over 5 hours and the  
pressure is reduced to 10 $\mu$ bar.

37. (Original) A lyophilized pharmaceutical composition  
prepared by the process of claim 27.

Claims 38-46. (Canceled)

47. (Previously Presented) The process of claim 27, wherein  
step b-i) is performed within 6 hours;  
step b-ii) is performed within 3 hours;  
step c-iii) is performed over 19 hours and at a  
pressure of 150 $\mu$ bar;  
step d-iv) is performed over 13 hours and at a  
pressure of 150 $\mu$ bar; and  
step e-v) is performed over 8 hours and at a  
pressure of 150 $\mu$ bar.

Claims 48-51. (Canceled)

52. (Previously Presented) A packaged pharmaceutical  
composition comprised of:  
a packaging material; and  
the lyophilized pharmaceutical composition of claim 37.